

**CRF Errors Corrected by the STIC Systems Branch**

Serial Number: 8/333,680

CRF Processing Date: 12/05/94  
 Edited by: W  
 Verified by: MM (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☒ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Inserted a space between the last nucleic designator and the nucleic number for sequences: \_\_\_\_\_
- ☐ Deleted page numbers in the text of the sequence listing, which is considered invalid text.
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted non-ASCII "garbage" at the end of files, and other invalid text, such as a secretary's initials.
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Other: \_\_\_\_\_

**ENTERED**

**\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.**

8/01/93

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/333,680DATE: 12/05/94  
TIME: 15:50:13

INPUT SET: S1373.raw

This Raw Listing contains the General  
Information Section and up to the first 5 pages.

## SEQUENCE LISTING

ENTERED

## (1) General Information:

(i) APPLICANT: Wang, Qing  
Finer, Mitchell H  
Jia, Xiao-Chi

(ii) TITLE OF INVENTION: NOVEL PACKAGING CELL LINES,  
ADENOVIRAL  
VECTORS AND RECOMBINANT ADENOVIRUSES AND METHODS

(iii) NUMBER OF SEQUENCES: 2

## (iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: CELL GENESYS, INC.  
(B) STREET: 322 Lakeside Drive  
(C) CITY: Foster City  
(D) STATE: California  
(E) COUNTRY: USA  
(F) ZIP: 94404

## (v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk  
(B) COMPUTER: IBM PC compatible  
(C) OPERATING SYSTEM: PC-DOS/MS-DOS  
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25

## (vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US  
(B) FILING DATE:  
(C) CLASSIFICATION:

## (viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Krupen, Karen  
(B) REGISTRATION NUMBER: 34,647

## (ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (415) 358-9600  
(B) TELEFAX: (415) 349-7392

## (2) INFORMATION FOR SEQ ID NO:1:

## (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 31 base pairs

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/333,680DATE: 12/05/94  
TIME: 15:50:17

INPUT SET: S1373.raw

47 (B) TYPE: nucleic acid  
48 (C) STRANDEDNESS: single  
49 (D) TOPOLOGY: linear  
50

51 (ii) MOLECULE TYPE: cDNA  
52  
53  
54

55 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:  
56

57 GCGCAAGCTT CGGGAGTGGG AGATAAGGCT C  
58

31

59 (2) INFORMATION FOR SEQ ID NO:2:  
60

61 (i) SEQUENCE CHARACTERISTICS:  
62 (A) LENGTH: 31 base pairs  
63 (B) TYPE: nucleic acid  
64 (C) STRANDEDNESS: single  
65 (D) TOPOLOGY: linear  
66

67 (ii) MOLECULE TYPE: cDNA  
68  
69  
70

71 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:  
72

73 GGCCCTCTAGA AGTTCACCTG CCCTGATGAC A  
74

31

INPUT SET: S1373.raw

## \*\*\*\*\* PREVIOUSLY ERRORED SEQUENCES - EDITED \*\*\*\*\*

43 (2) INFORMATION FOR SEQ ID NO:1:

44

45 (i) SEQUENCE CHARACTERISTICS:

46 (A) LENGTH: 31 base pairs

47 (B) TYPE: nucleic acid

48 (C) STRANDEDNESS: single

49 (D) TOPOLOGY: linear

50

51 (ii) MOLECULE TYPE: cDNA

52

53

54

55 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

56

57 GCGCAAGCTT CGGGAGTGGG AGATAAGGCT C

31

58

59 (2) INFORMATION FOR SEQ ID NO:2:

60

61 (i) SEQUENCE CHARACTERISTICS:

62 (A) LENGTH: 31 base pairs

63 (B) TYPE: nucleic acid

64 (C) STRANDEDNESS: single

65 (D) TOPOLOGY: linear

66

67 (ii) MOLECULE TYPE: cDNA

68

69

70

71 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

72

73 GGCCTCTAGA AGTTCACCTG CCCTGATGAC A

31

74

PAGE: 1

**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/333,680**

DATE: 12/05/94  
TIME: 15:50:23

*INPUT SET: S1373.raw*


Line	Error	Original Text
30	Wrong application Serial Number	(A) APPLICATION NUMBER: US

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/333,680DATE: 12/05/94  
TIME: 12:43:51

INPUT SET: S1373.raw

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

## SEQUENCE LISTING

1  
2  
3 (1) General Information:   
4  
5 (i) APPLICANT: Wang, Qing  
6                      Finer, Mitchell H  
7                      Jia, Xiao-Chi  
8  
9 (ii) TITLE OF INVENTION: NOVEL PACKAGING CELL LINES,  
10 ADENOVIRAL  
11                      VECTORS AND RECOMBINANT ADENOVIRUSES AND METHODS  
12  
13 (iii) NUMBER OF SEQUENCES: 2  
14  
15 (iv) CORRESPONDENCE ADDRESS:  
16                      (A) ADDRESSEE: CELL GENESYS, INC.  
17                      (B) STREET: 322 Lakeside Drive  
18                      (C) CITY: Foster City  
19                      (D) STATE: California  
20                      (E) COUNTRY: USA  
21                      (F) ZIP: 94404  
22  
23 (v) COMPUTER READABLE FORM:  
24                      (A) MEDIUM TYPE: Floppy disk  
25                      (B) COMPUTER: IBM PC compatible  
26                      (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
27                      (D) SOFTWARE: PatentIn Release #1.0, Version #1.25  
28  
29 (vi) CURRENT APPLICATION DATA:  
30                      (A) APPLICATION NUMBER: US  
31                      (B) FILING DATE:  
32                      (C) CLASSIFICATION:  
33  
34 (viii) ATTORNEY/AGENT INFORMATION:  
35                      (A) NAME: Krupen, Karen  
36                      (B) REGISTRATION NUMBER: 34,647  
37  
38 (ix) TELECOMMUNICATION INFORMATION:  
39                      (A) TELEPHONE: (415) 358-9600  
40                      (B) TELEFAX: (415) 349-7392  
41  
42

--&gt; ERROR SEQUENCES FOLLOW:

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/333,680DATE: 12/05/94  
TIME: 12:43:55

INPUT SET: S1373.raw

43 (2) INFORMATION FOR SEQ ID NO:1:

44

45 (i) SEQUENCE CHARACTERISTICS:

--&gt; 46 (A) LENGTH: 31 base pairs

47 (B) TYPE: nucleic acid

48 (C) STRANDEDNESS: single

49 (D) TOPOLOGY: linear

50

51 (ii) MOLECULE TYPE: cDNA

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53

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55 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

56

57 GCGCAAGCTT CGGGAGTGGG AGATAAGGCT C

58 31

59

60 (2) INFORMATION FOR SEQ ID NO:2:

61

62 (i) SEQUENCE CHARACTERISTICS:

--&gt; 63 (A) LENGTH: 31 base pairs

64 (B) TYPE: nucleic acid

65 (C) STRANDEDNESS: single

66 (D) TOPOLOGY: linear

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68 (ii) MOLECULE TYPE: cDNA

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71

72 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

73

74 GGCCTCTAGA AGTTCACTTG CCCTGATGAC A

75 31

76

PAGE: 1

**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/333,680**

DATE: 12/05/94  
TIME: 12:43:59

**INPUT SET: S1373.raw**

Line	Error	Original Text
30	Wrong application Serial Number	(A) APPLICATION NUMBER: US
46	Entered (31) and Calc. Seq. Length (0) differ	(A) LENGTH: 31 base pairs
63	Entered (31) and Calc. Seq. Length (0) differ	(A) LENGTH: 31 base pairs